

Prevalence of Tick Infestations and Tick-Borne Diseases in Cattle in Cameroon

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Abstract

Ticks induce huge production and economic losses in the livestock industry and create serious environmental, animal and human health problems. The study was carried out to characterize tick species and determine the prevalence of tick infestation and tick-borne diseases in cattle in Cameroon. Tick and blood samples were collected from a total of 742 animals and analyzed to determine the type of tick and haemoparasites using standard procedures. Overall, four tick species namely Amblyomma variegatum (75.09%), Rhipicephalus microplus (19.43%), Rhipicephalus decoloratus (0.88%) and Hyalomma marginatum (0.18%) and six blood disease agents including Anapalasma marginale (11.29%), Ehrlichia ruminantium (3.52%), Babesia bovis (1.32%), Babesia major (0.44%), Anaplasma phagocytophilum (0.29%) and Dermatophilus congolensis (3.37%). Various co-infections were recorded and the predominant associations were Amblyomma variegatum-Rhipicephalus microplus (4.06%) and Amblyomma variegatum-Hyalomma marginatum (0.36%); Anaplasma-Ehrlichia (12%), Anaplasma-Dematophylus (8%), Babesia-Anaplasma-Ehrlichia (14%) and Babesia-Anaplasma-Ehrlichia-Dermatophylus (5%). Breed, sex, age and locality significantly influenced the rate of tick infestation while locality, breed and age significantly influenced the detection of blood disease agents in the study.

Keywords

Prevalence, Tick, Blood Disease, Cattle, Cameroon

1. Introduction

The livestock sector accounts for over 40% of global agricultural production and contributes to the livelihoods and food security of nearly one billion people worldwide [1]. Livestock farming plays an important role in rural Africa [2] [3], and cattle are the main savings species for smallholders. Cattle are the fastest growing sector in the agricultural economy for income growth, infrastructure and technology [4]. The cattle population in Cameroon is estimated at 8,761,385 heads, and a meat beef production of 110,000 to 145,000 tons per year is equivalent to about 54% of meat products [5].

Ticks and tick-borne diseases are major problems for human, animal and environmental health with enormous negative economic impacts and financial losses of more than \$18.7 billion per year [6] [7] on beef production. Ticks are vectors of zoonoses of major public health importance [8] while tick-borne animal piroplasmosis and cowdriosis are major pathological and economic constraints on livestock development in sub-Saharan Africa [9].

Several control methods for ticks and tick-borne diseases (TBD) have been evaluated including the use of chemical products and acaricides against the vectors which pose a huge risk of drug resistance and pollution [10] [11] [12] [13]. Though the high cost of tick and tick-borne disease prevents expansion in cattle production [14] [15], vaccination is an alternative control tool in regions where there are no cross-border movements of cattle. Endogenous methods for tick control [8] [16] and other alternatives such as genetics geared towards using biological resistance mechanisms of certain species or breeds in livestock farming activities in tropical environments [17] [18] have been described.

The direct and indirect impacts of tick infestations of cattle are numerous including decreased milk productivity and carcass yield, low growth rate, the transmission of pathogens (eg. hemoparasites), dermatitis which may become infected by bacteria, skin coat depreciation making it less marketable and increase the cost of production due to tick control [19] [20] [21]. In the central African subregion including Cameroon, attention has focused on infectious and contagious diseases of animals while has neglected vector borne diseases especially tick-borne diseases [21] [22] [23].

Though there are increasing urban and peri-urban livestock production systems in most of Africa, there is a dearth of information on the level of awareness of cattle breeders on the hazards caused by ticks on husbandry systems and ectoparasite status of cattle in urban areas of Cameroon. In view of the medical and veterinary importance of ticks in the livestock sector, this study was carried out to determine the prevalence of tick infestation and ticks-borne diseases in cattle in Cameroon.

2. Materials and Methods

2.1. Study Location and Management of Animals

The study was carried out to determine the prevalence and intensity of tick in-

festation and haemoparasites in cattle within and around the SODEPA ranches (Livestock Development Corporation) in the Administrative Divisions of Faro & Deo (Adamawa region), Lom & Djerem (East region) in the Guinean high savannah agro-ecological zone and Donga-Mantung (North West region) in the western highland agro-ecological zone of Cameroon (Figure 1). The geographical and animal husbandry situations of the study regions are presented in Table 1 and Table 2.

The selection of cattle herds in the SODEPA Faro, Dumbo and Ndokayo ranches and those of willing local farmers in the environs of these SODEPA ranches during the study period, May to October 2021, was done using a random number generation technique from the list herds in the area [26]. Briefly, in each ranch, two large herds from among herds in the female camps to enable sufficient sampling of adult and young animals, and two herds from among herds in the adult male camps were included in the study. All individual animals in the chosen herds were sampled. Overall, tick and blood samples were collected from 742 cattle (409 Goudali and 333 Simgoud breeds; 337 males and 405 females; 90 young, 264 juveniles and 388 adults) corresponding to 268 cattle in Faro & Deo, Adamawa, 186 cattle in Lom & Djerem, East and 288 cattle in Donga-Mantung, North-West.

Following rigorous visual examination of tick on the preferred sites of the skin of the animals (anogenital, udder, perineum, abdomen, armpit and neck) [27], all the ticks encountered on the animal's body was harvested, by gently tapping at the rostrum and traction using a pair of forceps. The harvested ticks were stored in individual 5 ml labelled (identifying the sampled animal, anatomical region, and date of harvest) tubes containing 70% ethanol and 30% glycerol for analysis within 5 - 7 days in the parasitological laboratory of the *Mission Spéciale d'Eradication des Glossines (MSEG)* in *N'Gaoundéré*. Systemic phenotypic identification of the tick species was done using a binocular microscope and guided by reference tick images as previously described [28] [29] [30] [31] [32].

Blood samples (5 ml) from each animal were collected by jugular vein puncture into EDTA tubes for haematological analysis (Haematocrit value) and detection of haemoparasites using standard procedures. Haematocrits were determined to assess the degree of anaemia while blood smeared slides were stained with MAY-GRÜNWALD GIEMSA and observed under a light microscope at x100 magnification to detect the presence of haemoparasites.

Agro-ecological zone	Region	Surface area (km²)	Number of inhabitants	Climate	Average rainfall (mm)	Average altitude (m)	Cartesian coordinates
Guinean high savannah	Adamawa	6761	441,716	Sudanian tropical	950	950	7°05'06.00"N 13°12'10.80"E
	East	109,002	835,642	subtropical	1477	650	3°31'0"N 15°3'0"E
Western highlands	North west	17,812	1,840,500	savanna	916,6	1550	6°26'02.40"N 10°24'00.00"E

Table 1. Geographical characteristics of the study regions.

Source: Djoufack, 2011 [24].

Agro-ecological zone	Region	Type of farming	Cattle	sheep	Goats	Pigs	Poultry	Horse, donkey and camel
Guinean high savannah	Adamawa	Agropastoral and ranching	1,388,755	106,486	87,441	2668	248,744	6608
	East	Agropastoral and ranching	121,143	59,258	79,318	38,779	931,438	1925
Western highlands	North west	Agropastoral and ranching	467,817	240,712	368,190	201,917	756,414	23,247

Table 2. Animal husbandry features of the study regions.

source: MINEPIA/DEPCS, 2020 [25].

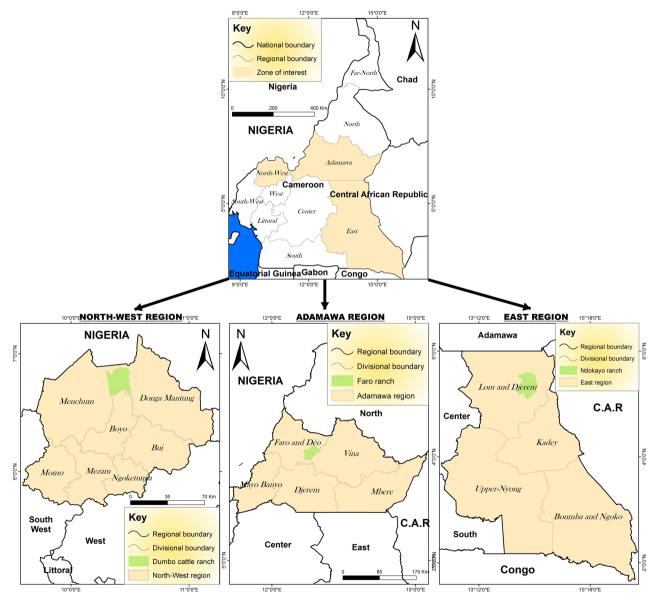


Figure 1. Map of Cameroon showing study regions and administrative divisions within the regions.

Information related to the breed, sex, age and body condition score of the sampled animals were noted. Estimation of ages was done by dental inspection

and examination of the horn rings for animals without teeth (especially old/adult females) while the breed of the animals was obtained as previously described [33] [34] [35]. The body condition score was done by assessing the general appearance and palpation of the lumbar region of the animal on a scale of 1 to 5 and further classed into 3 categories: 1 - 2 (poor), 3 (good) and 4 - 5 (very good) as previously described [36].

2.2. Ethical Consideration

Risk assessments of the project were performed by the researchers to avoid hazards to all persons involved in the project. Permission for the study was obtained from the required authorities in Cameroon including the MINEPIA delegations in Adamawa, East and Northwest and North Regions, SODEPA Cameroon (SODEPA Faro, Dumbo and Ndokayo ranches) and the Faculty of Agronomy and Agricultural Sciences of the University of Dschang, Cameroon. The purpose of the study was explained (with the assistance of local community leaders and trusted intermediaries) to cattle professionals with herds in the environs of the SODEPA ranches used in this study. Animals and farms were used in the study after informed consent was given by the owners.

2.3. Data Analysis

The data were analysed using SPSS 21.1 statistical software. Simple percentages were generated and Chi-square test was used to assess the association between factors and the odds-ratios determined for associated risk factors along 95% confidence intervals [26] [37]. The statistical significance was set at P < 0.05.

3. Results

3.1. Prevalence of Tick Infestation in Cattle in Cameroon

In this study, of the 742 cattle sampled 566 (76.28%, 95% CI: 73.00 - 79.56) animals were infested with ticks. Locality, age, breed and sex had significant effects (P < 0.05) on the prevalence of tick infestation in cattle in the study. Tick infestation was significantly higher in the Adamawa region (99.25%) than in the other 2 regions, in adult animals over 24 months of age (86.34%), in the Simgoud breed (79.87%) and in males (82.49%) (**Table 3**).

Microscopic examination of ticks taken from the cattle sampled revealed the presence of three genera (4 species) of ticks, including *Amblyomma variegatum* (75.09%), *Rhipicephalus microplus* (19.43%), *Rhipicephalus decoloratus* (0.88%) and *Hyalomma marginatum* (0.18%). There are also associations of ticks on the same animal. This was the case for *A. variegatum/R. microplus* (4.06%) and *A. variegatum/H. marginatum* (0.36%) (Table 4).

3.2. Prevalence of Tick-Borne Blood Diseases in Cattle in Cameroon

In this study, of the 742 cattle sampled 682 (91.91%, 95% CI) animals were in-

fected with tick-borne blood diseases. Locality, age, and breed had significant effects (P < 0.05) on the prevalence of tick-borne disease infection in cattle in the study while sex did not a significant effect (P > 0.05). Tick-borne disease infections were significantly higher in the East region (97.31%) than in the other 2 regions, in juvenile animals between 7 to 24 months of age (93.94%) and in the Simgoud breed (93.69%) (**Table 5**).

Factors	Variables	Number (positive)	Infestation rate % (95% CI)	СТ
	Adamawa	268 (266)	99.25	
Region	East	186 (109)	58.60	*
	Northwest	288 (191)	66.32	
	[0 - 6]	90 (52)	57.78	
Age class (month)]7 - 24]	264 (179)	67.80	*
	[24+[388 (335)	86.34	
D 1	Goudali	409 (300)	73.35	¥
Breed	Simgoud	333 (266)	79.87	2
C	Male	337 (278)	82.49	×
Sex	Female	405 (288)	71.11	88.60 * 66.32 * 97.78 * 77.80 * 78.634 * 79.87 * 12.49 * 111 *
	Total	742 (566)	76.28	

Table 3. Prevalence and risk factor for tick infestation of cattle in Cameroon.

CT: Contingency test, ns: non-significant, *: significant.

Table 4. Distributi	on of tick infested	d cattle according	to locality, bre	eed, age and sex in	Cameroon.

			Rhi	picephalus spp					Grand
Factors	Variables	Amblyomma variegatum	Rhipicephalus microplus	Rhipicephalus decoloratus	Total	•	•	A.variegatum/ H. marginatum	total number of infested cattle (%)
Total		425# (75.09)*	110 (19.43)	5 (0.88)	115 (20.31)	1 (0.18)	23 (4.06)	2 (0.36)	566 (100)
	Adamawa	253 (95.11)	1 (0.38)	0 (0)	1 (0.38)	0 (0)	11 (4.14)	1 (0.38)	266 (47)
Region	East	108 (99.08)	0 (0)	0 (0)	0 (0)	0 (0)	1 (0.92)	0 (0)	109 (19.26)
	Northwest	64 (33.51)	109 (57.07)	5 (2.62)	114 (59.69)	1 (0.52)	11 (5.76)	1 (0.52)	191 (33.75)
Age	[0 - 6]	40 (76.92)	11 (21.15)	0 (0)	11 (21.15)	0 (0)	1 (1.92)	0 (0)	52 (9.19)
class]7 - 24]	131 (73.18)	44 (24.58)	0 (0)	44 (24.58)	0 (0)	4 (2.23)	0 (0)	179 (31.63)
(month)	[24+[254 (75.82)	55 (16.42)	5 (1.49)	60 (19.91)	1 (0.30)	18 (5.37)	2 (0.60)	335 (59.19)
D	Goudali	228 (76)	63 (21)	0 (0)	63 (21)	0 (0)	7 (2.33)	2 (0.67)	300 (53)
Breed	Simgoud	197 (74.06)	47 (17.67)	5 (1.88)	52 (19.55)	1 (0.38)	16 (6.02)	0 (0)	266 (47)
	Male	219 (78.78)	37 (13.31)	1 (0.36)	38 (13.67)	0 (0)	19 (6.83)	2 (0.72)	278 (49.12)
Sex	Female	206 (71.53)	73 (25.35)	4 (1.39)	77 (26.74)	1 (0.35)	4 (1.39)	0 (0)	288 (50.88)

#: number cattle infested by tick, *: proportion (%) of infested cattle.

Microscopic examination of blood smears taken from the cattle sampled revealed the presence of three genera (5 species) of tick-borne blood diseases, including *Babesia bovis* (1.32%), *Babesia major* (0.44%), *Anaplasma marginale* (11.29%), *Anaplasma phagocytophylum* (0.29%) and *Ehrlichia ruminantium* (3.52%). There are also many associations of tick-borne diseases on the same animal. This represents the vast majority of tick-borne blood diseases (79.77%). We also identified *Dermatophilus congolensis* (3.37%), which is not a tick-borne blood disease but is strongly associated with the presence of *Amblyomma variegatum* (Table 6).

Factors	Variables	Number (positive)	Infection rate % (95% CI)	СТ
	Adamawa	268 (228)	85.07	
Region	East	186 (181)	97.31	*
	Northwest	288 (273)	94.79	
Age class (month)	[0 - 6]	90 (78)	86.67	
]7 - 24]	264 (248)	93.94	*
	[24+[388 (356)	91.75	
Dural	Goudali	409 (370)	90.46	*
Breed	Simgoud	333 (312)	93.69	
C	Male	337 (311)	92.28	
Sex	Female	405 (371)	91.60	ns
	Total	742 (682)	91.91	

Table 5. Prevalence and risk factor for tick-borne blood diseases infection of cattle in Cameroon.

CT: Contingency test, ns: non-significant, *: significant.

Table 6. Distribution of tick-borne blood diseases in cattle according to locality, breed, age and sex in Cameroon.

	Babesia spp			Апарі	Anaplasma spp			Dermatonhillus		Grand total number
Variables	Babesia bovis	Babesia major	Total	Anaplasma phagocytophylum	Anaplasma marginale	Total		congolensis	Polymorphism	of infested cattle (%)
	9# (1.32)*	3 (0.44)	12 (1.76)	2 (0.29)	77 (11.29)	79 (11.58)	24 (3.52)	23 (3.37)	544 (79.77)	682 (100.00)
Adamawa	9 (3.95)	2 (0.88)	11 (4.83)	2 (0.88)	15 (6.58)	17 (7.46)	8 (3.51)	18 (7.89)	174 (76.32)	228 (33.43)
East	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	15 (8.29)	15 (8.29)	4 (2.21)	0 (0.00)	162 (89.50)	181 (26.54)
Northwest	0 (0.00)	1 (0.37)	1 (0.37)	0 (0.00)	47 (17.22)	47 (17.22)	11 (4.03)	6 (2.20)	208 (76.19)	273 (40.03)
[0 - 6]	0 (0.00)	0 (0.00)	0 (0.00)	0 (0.00)	8 (10.26)	8 (10.26)	0 (0.00)	2 (2.56)	68 (87.18)	78 (11.44)
]7 - 24]	1 (0.40)	2 (0.81)	3 (1.21)	0 (0.00)	11 (4.44)	11 (4.44)	4 (1.61)	9 (3.63)	221 (89.11)	248 (36.36)
[24+[8 (2.25)	1 (0.28)	9 (2.53)	2 (0.56)	58 (16.29)	60 (16.85)	19 (5.34)	13 (3.65)	255 (71.63)	356 (52.20)
Goudali	7 (1.89)	0 (0.00)	7 (1.89)	1 (0.27)	40 (10.81)	41 (11.08)	17 (4.59)	20 (5.41)	285 (77.03)	370 (54.25)
Simgoud	2 (0.64)	3 (0.96)	5 (1.6)	1 (0.32)	37 (11.86)	38 (12.18)	6 (1.92)	4 (1.28)	259 (83.01)	312 (45.75)
Male	3 (0.96)	2 (0.64)	5 (1.6)	1 (0.32)	36 (11.58)	37 (11.90)	10 (3.22)	13 (4.18)	246 (79.10)	311 (45.60)
Female	6 (1.62)	1 (0.27)	7 (1.89)	1 (0.27)	41 (11.05)	42 (11.32)	13 (3.50)	11 (2.96)	298 (80.32)	371 (54.40)
	Adamawa East Northwest [0 - 6]]7 - 24] [24+[Goudali Simgoud Male	Babesia Babesia bvvis 9°(1.32)* Adamawa 9°(0.000) East 000000 Fast 010000 10000 10101 10102 10103 10241 101040 10105 10106 10107 10108 10109	Babesia Babesia Pariables Babesia major 9* (1.32)* 3 (0.44) Adamawa 9 (3.95) 2 (0.88) East 0 (0.00) 0 (0.00) Rorthwest 0 (0.00) 1 (0.37) [0 - 6] 0 (0.00) 0 (0.00)]7 - 24] 1 (0.40) 2 (0.81) [24+[8 (2.25) 1 (0.28) Goudali 7 (1.89) 0 (0.00) Simgoud 2 (0.64) 3 (0.96)	Variables Babesia bovis Babesia major Total 9* (1.32)* 3 (0.44) 12 (1.76) Adamawa 9 (3.95) 2 (0.88) 11 (4.83) East 0 (0.00) 0 (0.00) 0 (0.00) Northwest 0 (0.00) 1 (0.37) 1 (0.37) [0 - 6] 0 (0.00) 0 (0.00) 0 (0.00)]7 - 24] 1 (0.40) 2 (0.81) 3 (1.21) [24+[8 (2.25) 1 (0.28) 9 (2.53) Goudali 7 (1.89) 0 (0.00) 7 (1.89) Simgoud 2 (0.64) 3 (0.96) 5 (1.6)	Variables Babesia bovis Babesia major Total Anaplasma phagocytophylum 9* (1.32)* 3 (0.44) 12 (1.76) 2 (0.29) Adamawa 9 (3.95) 2 (0.88) 11 (4.83) 2 (0.88) East 0 (0.00) 0 (0.00) 0 (0.00) 0 (0.00) Northwest 0 (0.00) 1 (0.37) 1 (0.37) 0 (0.00) [0 - 6] 0 (0.00) 0 (0.00) 0 (0.00) 0 (0.00) [7 - 24] 1 (0.40) 2 (0.81) 3 (1.21) 0 (0.00) [24+[8 (2.25) 1 (0.28) 9 (2.53) 2 (0.56) Goudali 7 (1.89) 0 (0.00) 7 (1.89) 1 (0.27) Simgoud 2 (0.64) 3 (0.96) 5 (1.6) 1 (0.32)	Variables Babesia bovis Babesia major Total Anaplasma phagocytophylum Anaplasma marginale 9* (1.32)* 3 (0.44) 12 (1.76) 2 (0.29) 77 (11.29) Adamawa 9 (3.95) 2 (0.88) 11 (4.83) 2 (0.88) 15 (6.58) East 0 (0.00) 0 (0.00) 0 (0.00) 0 (0.00) 15 (8.29) Northwest 0 (0.00) 1 (0.37) 1 (0.37) 0 (0.00) 47 (17.22) [0 - 6] 0 (0.00) 0 (0.00) 0 (0.00) 0 (0.00) 47 (17.22) [0 - 6] 0 (0.00) 0 (0.00) 0 (0.00) 0 (0.00) 11 (4.44) [24+[8 (2.25) 1 (0.28) 9 (2.53) 2 (0.56) 58 (16.29) Goudali 7 (1.89) 0 (0.00) 7 (1.89) 1 (0.27) 40 (10.81) Simgoud 2 (0.64) 3 (0.96) 5 (1.6) 1 (0.32) 36 (11.58)	Variables Babesia bovis Babesia major Total Anaplasma phagocytophylum Anaplasma marginale Total 9* (1.32)* 3 (0.44) 12 (1.76) 2 (0.29) 77 (11.29) 79 (11.58) Adamawa 9 (3.95) 2 (0.88) 11 (4.83) 2 (0.29) 77 (11.29) 79 (11.58) Adamawa 9 (3.95) 2 (0.88) 11 (4.83) 2 (0.88) 15 (6.58) 17 (7.46) East 0 (0.00) 0 (0.00) 0 (0.00) 0 (0.00) 15 (8.29) 15 (8.29) Northwest 0 (0.00) 1 (0.37) 1 (0.37) 0 (0.00) 47 (17.22) 47 (17.22) [0 - 6] 0 (0.00) 0 (0.00) 0 (0.00) 0 (0.00) 8 (10.26) 8 (10.26)]7 - 24] 1 (0.40) 2 (0.81) 3 (1.21) 0 (0.00) 11 (4.44) 11 (4.44) [24+[8 (2.25) 1 (0.28) 9 (2.53) 2 (0.56) 58 (16.29) 60 (16.85) Goudali 7 (1.89) 0 (0.00) 7 (1.89) 1 (0.27) 40 (10.81) 41 (11.08)	VariablesBabesia bovisBabesia majorTotal rtalAnaplasma phagocytophylumAnaplasma marginaleAnaplasma ruminantium9* (1.32)*3 (0.44)12 (1.76)2 (0.29)77 (11.29)79 (11.58)24 (3.52)Adamawa9 (3.95)2 (0.88)11 (4.83)2 (0.88)15 (6.58)17 (7.46)8 (3.51)East0 (0.00)0 (0.00)0 (0.00)0 (0.00)15 (8.29)15 (8.29)4 (2.21)Northwest0 (0.00)1 (0.37)1 (0.37)0 (0.00)47 (17.22)11 (4.03)[0 - 6]0 (0.00)0 (0.00)0 (0.00)0 (0.00)8 (10.26)8 (10.26)0 (0.00) $[7 - 24]$ 1 (0.40)2 (0.81)3 (1.21)0 (0.00)11 (4.44)11 (4.44)4 (1.61) $[24+[$ 8 (2.25)1 (0.28)9 (2.53)2 (0.56)58 (16.29)60 (16.85)19 (5.34)Goudali7 (1.89)0 (0.00)7 (1.89)1 (0.27)40 (10.81)41 (11.08)17 (4.59)Simgoud2 (0.64)3 (0.96)5 (1.6)1 (0.32)36 (11.58)37 (11.90)10 (3.22)	VariablesBabesia bovisBabesia majorTotal rotalAnaplasma phagocytophylumAnaplasma marginaleTotalEhrlichia ruminantiumDermatophillus congolensis9* (1.32)*3 (0.44)12 (1.76)2 (0.29)77 (11.29)79 (11.58)24 (3.52)23 (3.37)Adamawa9 (3.95)2 (0.88)11 (4.83)2 (0.88)15 (6.58)17 (7.46)8 (3.51)18 (7.89)East0 (0.00)0 (0.00)0 (0.00)0 (0.00)15 (8.29)15 (8.29)4 (2.21)0 (0.00)Northwest0 (0.00)1 (0.37)1 (0.37)0 (0.00)47 (17.22)47 (17.22)11 (4.03)6 (2.20)[0 - 6]0 (0.00)0 (0.00)0 (0.00)0 (0.00)8 (10.26)8 (10.26)0 (0.00)2 (2.56) $[7 - 24]$ 1 (0.40)2 (0.81)3 (1.21)0 (0.00)11 (4.44)11 (4.44)4 (1.61)9 (3.63) $[24+[$ 8 (2.25)1 (0.28)9 (2.53)2 (0.56)58 (16.29)60 (16.85)19 (5.34)13 (3.65)Goudali7 (1.89)0 (0.00)7 (1.89)1 (0.27)40 (10.81)41 (11.08)17 (4.59)20 (5.41)Simgoud2 (0.64)3 (0.96)5 (1.6)1 (0.32)37 (11.86)38 (12.18)6 (1.92)4 (1.28)Male3 (0.96)2 (0.64)5 (1.6)1 (0.32)36 (11.58)37 (11.90)10 (3.22)13 (4.18)	Variables Babesia bovis Babesia major Total (1, 4, 8) Anaglasma phagocytophylum Anaglasma marginale Total Ehrlichia ruminantium Dermatophillus congolensis Polymorphism 9 ⁴ (1.32)* 3 (0.44) 12 (1.76) 2 (0.29) 77 (11.29) 79 (11.58) 24 (3.52) 23 (3.37) 544 (79.77) Adamawa 9 (3.95) 2 (0.88) 11 (4.83) 2 (0.88) 15 (6.58) 17 (7.46) 8 (3.51) 18 (7.89) 174 (76.32) East 0 (0.00) 0 (0.00) 0 (0.00) 0 (0.00) 160.289.50 16 (2.20) 208 (76.19) [0 - 6] 0 (0.00) 1 (0.37) 1 (0.37) 0 (0.00) 47 (17.22) 47 (17.22) 11 (4.03) 6 (2.20) 208 (76.19) [0 - 6] 0 (0.00) 1 (0.37) 1 (0.37) 0 (0.00) 8 (10.26) 8 (10.26) 0 (0.00) 2 (2.56) 68 (87.18) [7 - 24] 1 (0.40) 2 (0.81) 3 (1.21) 0 (0.00) 11 (4.44) 11 (4.44) 4 (1.61) 9 (3.63) 221 (89.11) [24+[] 8 (2.25)

#: number of cattle infected by haemoparasites, *: proportion (%) of infected cattle.

4. Discussion

The prevalence of ticks and TBD has been widely documented in tropical environments. This aspect is addressed in several ways by various methods, namely blood sampling and germ detection or molecular analysis of the ticks themselves. The results obtained on prevalence as a function of locality significantly corroborate those of [38], who argue that the prevalence of each tick species in its geographical distribution is conditioned by biomes and specific environmental conditions such as temperature. It is normal that, depending on the environmental conditions, there should be various adaptations of species.

4.1. Amblyomma variegatum

The results obtained from the frequency of this tick in the different regions of 75.09% show the abundance of this tick and are similar to those obtained from 46.3% to 73.8% [39]-[45]. These same authors found that it is practically the most widespread in several African countries. However, [46] found that the genus *Rhipicephalus* (80.60%) was more important than *Amblyomma* (19.31%). This tick thrives in the warmer regions of Africa. We note that it is less significant in the north-west, which is a mountainous area with low average annual temperatures compared with the other two regions.

4.2. Rhipicephalus microplus

The results obtained for this tick are 19.43% overall. However, it is very prevalent in the North West region, where 57.07% of animals are infested with this species, in contrast to the other two regions, where it was only mentioned very rarely. This prevalence is not far from the results obtained by numerous authors in several countries, including Cameroon [41] [43], which vary from 15.6% to 24.11%. This confirms the presence of this highly invasive species in Cameroon [47].

4.3. Hyalomma marginatum

The results obtained from the presence of this tick, concentrated in the north-west with a very low prevalence of around 0.18%, are similar to those obtained from several studies which only mention its presence without mentioning a precise prevalence. These include [39] [48] [49]. However, other authors have given prevalence ranging from 1.8% to 13.8% [45] [50] [51].

4.4. Rhipicephalus decoloratus

The results we obtained (0.88%) were founded only in the north-west among adults. However, several studies have mentioned its presence (20.1% to 50.6%) in various regions of Africa [38] [44] [52] [53]. Our results are higher than obtained in north-west Cameroon (0.4%) by [39].

The results obtained in the different regions mentioning the presence of babesiosis, anaplasmosis, richetsiosis and dermatophilosis are caused by several germs sometimes associated with each other on the same animal.

4.5. Babesiosis

We encountered two germs, *Babesia bovis* (1.3%) and *Babesia major* (0.44%), especially in the East region. The results are not very far from those obtained in small ruminants (2.9%) in the Adamawa region [41]. This disease is usually mentioned in the records of the veterinary services in all regions where cattle are reared.

4.6. Anaplasmosis

We encountered two germs, *Anaplasma marginale* (11.29%) and *Anaplasma phagocytophilum* (0.29%), respectively in the three regions for the former and only in Adamawa for the latter. The prevalence of *Anaplasma marginale* is similar to the results obtained for small ruminants (9.1%) in Adamawa [41]. This species is the most pathogenic of its genus [54].

4.7. Ehrlichia ruminantium

The results obtained for the prevalence of around 3.52% are different from those obtained in a study carried out in Benin using molecular analysis of ticks, which revealed the germ with an overall prevalence in ticks of 10.8% [49] and 28.4% [55]. This may be due to the different techniques used, as we worked on the animals and not on the ticks. In addition, there is a correlation between the presence of this germ and the environment.

5. Conclusion

This study was carried out to determine the prevalence and intensity of tick infestation and haemoparasites in cattle within and around the SODEPA ranches in the Guinean high savannah agro-ecological zone and in the western highland agro-ecological zone of Cameroon reveals the presence of three ticks genus (4 species): Amlyomma variegatum, Rhipicephalus microplus, Hyalomma marginatum and Rhipicephalus decoloratus. The following tick-borne diseases were also identified: Babesia bovis, Babesia major, Anaplasma marginale, Anaplasma phagocytophylum and Ehrlichia ruminantium. We also identified Dermatophilus congolensis which is not a tick-borne disease but associated with Amblyomma presence. There is a significant dependence between the prevalence of ticks and locality, sex, breed and age. Similarly, there was a dependence on the prevalence of tick-borne diseases and locality, breed and age. On the other hand, there was no dependence between tick-borne diseases and the sex of the animal. The perspective of this study is to examine the implementation of a tick and haemoparasite control mechanism that is both animal welfare-friendly and environmentally friendly.

Authors' Contributions

HH conceptualised the study under the direction of FM, and he collected data and wrote the first draft of the manuscript. AFE contributed to the final manu-

script. FM, RA, NAJ and BM supervised HH.

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Conflicts of Interest

The authors declare no conflicts of interest regarding the publication of this paper.

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